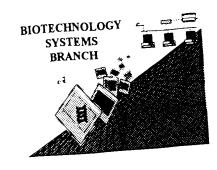
RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/83/,047ASource: 9/30/301Date Processed by STIC:

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE

TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST 25

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2Kcompliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

Raw Sequence Listing Error Summary

	20/021-411
	SUCCESTED CORRECTION SERIAL NUMBER: 09/83/,047A
TEROP DETECTED	SUGGESTED CORRECTION SERIAL NUMBERS PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE INSERTED BY P
ERROR DETECTION	THE ASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE
ATTN: NEW RULES CASES:	PLEASE DISREGARD ENGLISH ADVANCE The number/text at the end of each line "wrapped" down to the next line. This may occur if your file The number/text at the end of each line "wrapped" down to the next line. This may occur if your file The number/text at the end of each line "wrapped" down to the next line. This may occur if your file The number/text at the end of each line "wrapped" down to the next line. This may occur if your file The number/text at the end of each line "wrapped" down to the next line. This may occur if your file The number/text at the end of each line "wrapped" down to the next line. This may occur if your file The number/text at the end of each line "wrapped" down to the next line. This may occur if your file The number/text at the end of each line "wrapped" down to the next line. This may occur if your file The number/text at the end of each line "wrapped" down to the next line. This may occur if your file The number/text at the end of each line "wrapped" down to the next line. This may occur if your file The number/text at the end of each line "wrapped" down to the next line. This may occur if your file The number/text at the end of each line "wrapped" down to the next line. This may occur if your file The number/text at the end of each line "wrapped" down to the next line. This may occur if your file The number of the number of the number of the next line. This may occur if your file The number of
1Wrapped Nucleics	The number/text at the end of each line "wrapped" down to the next line. This may occur if you was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will was retrieved in a word processor after creating it.
	was retrieved in a word process
	prevent "wrapping." The rules require that a line not exceed 72 characters in length. This includes white spaces.
2Invalid Line Length	The rules require that a line not exceed 72 characters in rengine
	The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; The numbering under each 5th amino acid is misaligned.
3Misaligned Amino Numbering	The numbering under each) annul acts to
	The numbering characters, instead. Use space characters, instead. The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please The submitted file was not saved in ASCII text.
4Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by
	ensure your subsequent sales.
	Yea's representing more than one residue.
5Variable Length.	Sequence(s) contain n's or Xaa's representing more than one residue. Per sequence teach nor Xaa can only represent a single residue. Please present the maximum number of each each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
	each n or Ana car only serious in the <220>-<223> section management and indicate in the <220>-<223> section
	residue having variable length and indicate more residue having variable length and indicate more residue having variable length and indicate more residue having from amino acid A "bug" in Patentln version 2.0 has caused the <220>-<223> section to be missing from amino acid A "bug" in Patentln version 2.0 has caused the <220>-<223> section to secti
6PatentIn 2.0 "bug"	A "bug" in Patentin version 2.0 has caused the would automatically generate this section from the
	A "bug" in Patentln version 2.0 has caused the <220>-<223> section to be missing from animo and the sequences(s) Normally, Patentln would automatically generate this section from the sequences(s) previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> sections to previously coded nucleic acid sequence. This applies to the mandatory <220>-<223> sections for
	previously coded interest and sequence. This applies to the manuality
7Skipped Sequences (OLD RULES) 8Skipped Sequences (NEW RULES)	Artificial or Unknown sequences.
	Is intentional please insert the following times to
	Sequence(s) missing. It intentions are sequence(s) missing. It intentions are sequence(s) missing. It intentions are sequence (i) NFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (ii) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
	" anoticing DESCRIPTION DESCRIPTION OF A STATE OF THE STA
	(xi) SEQUENCE BEST This sequence is intentionally skipped This sequence is intentionally skipped sequences.
	OF SEOUENCES:" response to Bretade at the second of the se
	Please also adjust the "(ii) NUMBER Of SEQUENCE. Sequence(s) missing. If Intentional, please insert the following lines for each skipped sequence.
	Sequence(s) missing. If Intentional, please insert the room
	(210) sequence id number
	<400> sequence id number
1	000
	Use of n's and/or Xaa's have been detected in the Sequence Listing. Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>.<223> is MANDATORY if n's or Xaa's are present. Per 1.823 of Sequence Rules, use of <200>.<223> is MANDATORY if n's or Xaa represents.
9 Use of n's or Xaa's (NEW RULES)	Per 1.823 of Sequence Rules, location of n or Xaa, and which resident
(NEW ROLLS)	In <220> 10 <225 section F
	In <220> to <223> section, please explaints Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or Scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or scientific name (Genus/species).
10Invalid <213> Response	ecientific name (Genus/species). <220>-<223> section is require
	is Artificial Sequence
11Use of <220>	is Artificial Sequence missing the <220> "Feature" and associated numeric identifiers and responses. Sequence(s)
	Use of <220> to <223> is MANDATORY if <213> "Organism response is Vision." "Unknown." Please explain source of genetic material in <220> to <223> section. "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
	(See "Federal Register," 06/01/1998, Vol. 63, No. 13 4, Fr
12PatentIn 2.0 "bug"	Diek" function of Patentin version 2.0. This feeted on raw sequence
	Please do not use "Copy to Disk" function of Patentin version 2.0. This causes a corresponding to manual manual means to copy file to floppy disk.
	Please do not use "Copy to Disk Interest identifiers and responses (as indicated on the description of the resulting in missing mandatory numeric identifiers and responses (as indicated on the description of the resulting). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
	N is not used to represent
	n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent
13 Misuse of n	any value not specifically a nucleotide.
	any value not specially systems Branch - 08/21/2001

AMC/MH - Biotechnology Systems Branch - 08/21/2001



DATE: 09/20/2001 RAW SEQUENCE LISTING TIME: 08:44:21 PATENT APPLICATION: US/09/831,047A Input Set : A:\WST93AUSA.txt Output Set: N:\CRF3\09202001\1831047A.raw 4 <110> APPLICANT: The Wistar Institute of Anatomy and Biology The Trustees of the University of Pennsylvania Blaszczyk-Thurin, Magdalena 9 <120> TITLE OF INVENTION: Compositions and Methods For Treatment of Cancer 11 <130> FILE REFERENCE: WST93PCT C--> 13 <140> CURRENT APPLICATION NUMBER: US/09/831,047A C--> 14 <141> CURRENT FILING DATE: 2001-06-30 16 <150> PRIOR APPLICATION NUMBER: 60/107,478 17 <151> PRIOR FILING DATE: 1998-11-06 19 <160> NUMBER OF SEQ ID NOS: 121 21 <170> SOFTWARE: PatentIn Ver. 2.0 23 <210> SEQ ID NO: 1 24 <211> LENGTH: 12 25 <212> TYPE: PRT 26 <213> ORGANISM: Artificial Sequence 29 <223> OTHER INFORMATION: Description of Artificial Sequence: peptido-mimetic of a Lewis antigen 32 <400> SEQUENCE: 1 33 Asp Leu Trp Asp Trp Val Val Gly Lys Pro Ala Gly 5 34 1 37 <210> SEQ ID NO: 2 38 <211> LENGTH: 12 39 <212> TYPE: PRT 40 <213> ORGANISM: Artificial Sequence 43 <223> OTHER INFORMATION: Description of Artificial Sequence: peptido-mimetic of a Lewis antigen 44 47 Asp Ala Trp Asp Trp Val Val Gly Lys Pro Ala Gly 48 1 51 <210> SEQ ID NO: 3 52 <211> LENGTH: 12 53 <212> TYPE: PRT 54 <213> ORGANISM: Artificial Sequence 57 <223> OTHER INFORMATION: Description of Artificial Sequence: peptido-mimetic of a Lewis antigen 61 Asp Asp Trp Asp Trp Val Val Gly Lys Pro Ala Gly 62 1 65 <210> SEQ ID NO: 4 66 <211> LENGTH: 12 67 <212> TYPE: PRT 68 <213> ORGANISM: Artificial Sequence 70 <220> FEATURE:





DATE: 09/20/2001 TIME: 08:44:21 RAW SEQUENCE LISTING PATENT APPLICATION: US/09/831,047A

Input Set : A:\WST93AUSA.txt

Output Set: N:\CRF3\09202001\I831047A.raw

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71 <223> OTHER INFORMATION: Description of Artificial Sequence:
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75 Asp Tyr Trp Asp Trp Val Val Gly Lys Pro Ala Gly
                     5
76 1
79 <210> SEQ ID NO: 5
80 <211> LENGTH: 12
81 <212> TYPE: PRT
82 <213> ORGANISM: Artificial Sequence
 85 <223> OTHER INFORMATION: Description of Artificial Sequence:
        peptido-mimetic of a Lewis antigen
 88 <400> SEQUENCE: 5
 89 Asp Glu Trp Asp Trp Val Val Gly Lys Pro Ala Gly
                      5
 90 1
 93 <210> SEQ ID NO: 6
 94 <211> LENGTH: 12
 95 <212> TYPE: PRT
 96 <213> ORGANISM: Artificial Sequence
  99 <223> OTHER INFORMATION: Description of Artificial Sequence:
  100 peptido-mimetic of a Lewis antigen
  102 <400> SEQUENCE: 6
  103 Asp Lys Trp Asp Trp Val Val Gly Lys Pro Ala Gly
                        5
  1.04 1
  107 <210> SEQ ID NO: 7
  108 <211> LENGTH: 12
  109 <212> TYPE: PRT
  110 <213> ORGANISM: Artificial Sequence
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   117 Asp Arg Trp Asp Trp Val Val Gly Lys Pro Ala Gly
   118 1
   121 <210> SEQ ID NO: 8
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   123 <212> TYPE: PRT
   124 <213> ORGANISM: Artificial Sequence
   127 <223> OTHER INFORMATION: Description of Artificial Sequence:
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    130 <400> SEQUENCE: 8
    131 Asp Ser Trp Asp Trp Val Val Gly Lys Pro Ala Gly
    132 1
    135 <210> SEQ ID NO: 9
    136 <211> LENGTH: 12
    137 <212> TYPE: PRT
     138 <213> ORGANISM: Artificial Sequence
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DATE: 09/20/2001 RAW SEQUENCE LISTING TIME: 08:44:21 PATENT APPLICATION: US/09/831,047A

Input Set : A:\WST93AUSA.txt

Output Set: N:\CRF3\09202001\I831047A.raw

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141 <223> OTHER INFORMATION: Description of Artificial Sequence:
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144 <400> SEQUENCE: 9
145 Asp Leu His Asp Trp Val Val Gly Lys Pro Ala Gly
                   5
146 1
149 <210> SEQ ID NO: 10
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151 <212> TYPE: PRT
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 158 <400> SEQUENCE: 10
 159 Asp Leu Tyr Asp Trp Val Val Gly Lys Pro Ala Gly
 160 1
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 164 <211> LENGTH: 12
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  172 <400> SEQUENCE: 11
  173 Asp Leu Phe Asp Trp Val Val Gly Lys Pro Ala Gly
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  174 1
  177 <210> SEQ ID NO: 12
  178 <211> LENGTH: 12
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  180 <213> ORGANISM: Artificial Sequence
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   186 <400> SEQUENCE: 12
   187 Asp Leu Met Asp Trp Val Val Gly Lys Pro Ala Gly
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   188 1
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   193 <212> TYPE: PRT
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    205 <210> SEQ ID NO: 14
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DATE: 09/20/2001 TIME: 08:44:21 RAW SEQUENCE LISTING PATENT APPLICATION: US/09/831,047A

Input Set : A:\WST93AUSA.txt

Output Set: N:\CRF3\09202001\1831047A.raw

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208 <213> ORGANISM: Artificial Sequence
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215 Asp Leu Glu Asp Trp Val Val Gly Lys Pro Ala Gly
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  235 <212> TYPE: PRT
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  239 <223> OTHER INFORMATION: Description of Artificial Sequence:
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  240
   244 1
   247 <210> SEQ ID NO: 17
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    257 Asp Leu Arg Asp Trp Val Val Gly Lys Pro Ala Gly
    258 1
    261 <210> SEQ ID NO: 18
    262 <211> LENGTH: 12
    263 <212> TYPE: PRT
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     272 1
     275 <210> SEQ ID NO: 19
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DATE: 09/20/2001 TIME: 08:44:21 RAW SEQUENCE LISTING PATENT APPLICATION: US/09/831,047A

Input Set : A:\WST93AUSA.txt

Output Set: N:\CRF3\09202001\I831047A.raw

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277 <212> TYPE: PRT
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281 <223> OTHER INFORMATION: Description of Artificial Sequence:
280 <220> FEATURE:
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285 Asp Leu Trp Glu Trp Val Val Gly Lys Pro Ala Gly
286 1
289 <210> SEQ ID NO: 20
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<210> SEQ ID NO 100

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<223> OTHER INFORMATION: Description of Artificial Sequence: <220> FEATURE:

peptido-mimetic of a Lewis antigen

<400> SEQUENCE: 100

Asn Leu Arg Pro Lys Tyr Ile (Xaa) Leu Asp Ala Asp

Del item You Erre, Summary Skeet

sand enn 12 Sejs. 111-112





DATE: 09/20/2001 TIME: 08:44:22 VERIFICATION SUMMARY PATENT APPLICATION: US/09/831,047A

Input Set : A:\WST93AUSA.txt

Output Set: N:\CRF3\09202001\I831047A.raw

L:13 M:270 C: Current Application Number differs, Replaced Application Number L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:1419 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:100 L:1419 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:100 L:1419 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:100 L:1573 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:111 L:1573 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:111 L:1573 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:111 L:1587 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:112 L:1587 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:112 L:1587 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:112